



Author index volume 335 (2005)

The issue number is given in front of the page numbers.

Aceto, L., W. Fokkink, A. Ingolfsdottir and Z. Ésik, Foreword	(2–3) 127–129
Aldini, A. and M. Bernardo, On the usability of process algebra: An architectural view	(2–3) 281–329
Apostolico, A. and R. Giancarlo, Foreword	(1) 1– 2
Baeten, J.C.M., A brief history of process algebra	(2–3) 131–146
Bafna, V., S. Istrail, G. Lancia and R. Rizzi, Polynomial and APX-hard cases of the individual haplotyping problem	(1) 109–125
Bergstra, J.A. and C.A. Middelburg, Process algebra for hybrid systems	(2–3) 215–280
Bernardo, M., <i>see</i> A. Aldini	(2–3) 281–329
Bruni, R., U. Montanari and V. Sassone, Observational congruences for dynamically reconfigurable tile systems	(2–3) 331–372
Chattaraj, A. and L. Parida, An inexact-suffix-tree-based algorithm for detecting extensible patterns	(1) 3– 14
Choy, C., J. Jansson, K. Sadakane and W.-K. Sung, Computing the maximum agreement of phylogenetic networks	(1) 93–107
Corradini, F. and W. Vogler, Measuring the performance of asynchronous systems with PAFAS	(2–3) 187–213
Dong, G., <i>see</i> L. Wang	(1) 15– 27
Ésik, Z., <i>see</i> L. Aceto	(2–3) 127–129
Fokkink, W., <i>see</i> L. Aceto	(2–3) 127–129
Giancarlo, R., <i>see</i> A. Apostolico	(1) 1– 2
Herescu, O.M., <i>see</i> C. Palamidessi	(2–3) 373–404
Ingolfsdottir, A., <i>see</i> L. Aceto	(2–3) 127–129
Istrail, S., <i>see</i> V. Bafna	(1) 109–125
Jansson, J., <i>see</i> C. Choy	(1) 93–107
Lancia, G., <i>see</i> V. Bafna	(1) 109–125
Li, J., <i>see</i> L. Wang	(1) 15– 27
Li, J., <i>see</i> Y. Xu	(1) 53– 66
Luttik, B. and V. van Oostrom, Decomposition orders—another generalisation of the fundamental theorem of arithmetic	(2–3) 147–186
Mauri, G. and G. Pavesi, Algorithms for pattern matching and discovery in RNA secondary structure	(1) 29– 51
Middelburg, C.A., <i>see</i> J.A. Bergstra	(2–3) 215–280
Montanari, U., <i>see</i> R. Bruni	(2–3) 331–372
Palamidessi, C. and O.M. Herescu, A randomized encoding of the π -calculus with mixed choice	(2–3) 373–404
Parida, L., <i>see</i> A. Chattaraj	(1) 3– 14
Pavesi, G., <i>see</i> G. Mauri	(1) 29– 51
Rizzi, R., <i>see</i> V. Bafna	(1) 109–125
Sadakane, K., <i>see</i> C. Choy	(1) 93–107
Sassone, V., <i>see</i> R. Bruni	(2–3) 331–372
Steyaert, J.-M., <i>see</i> J. Waldisphl	(1) 67– 92
Sung, W.-K., <i>see</i> C. Choy	(1) 93–107
van Oostrom, V., <i>see</i> B. Luttik	(2–3) 147–186
Vogler, W., <i>see</i> F. Corradini	(2–3) 187–213

- Waldispohl, J. and J.-M. Steyaert, Modeling and predicting all- α transmembrane proteins including helix–helix pairing (1) 67– 92
- Wang, L., H. Zhao, G. Dong and J. Li, On the complexity of finding emerging patterns (1) 15– 27
- Wang, L., *see* Y. Xu (1) 53– 66
- Xu, Y., L. Wang, H. Zhao and J. Li, Exact matching of RNA secondary structure patterns (1) 53– 66
- Zhao, H., *see* L. Wang (1) 15– 27
- Zhao, H., *see* Y. Xu (1) 53– 66